

**Go to:** General Description References Cross-references Sequence

## General Information

XP-002386149

Primary Accession # BQ055725

Accession # BQ055725

Entry Name EMBL:BQ055725

Molecule Type linear mRNA

Sequence Length 1060

Entry Division HUM

Entry Data Class EST

Sequence Version BQ055725.1

Creation Date 01-APR-2002

Modification Date 01-APR-2002

## Description

Description AGENCOURT\_6796360 NIH\_MGC\_99 Homo sapiens cDNA clone IMAGE:5807946 5', mRNA sequence.

Keywords EST,;

Organism Homo sapiens (human)

Organism Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia;  
Classification Eutheria; Euarchontoglires; Primates; Haplorrhini; Catarrhini; Hominidae; Homo.

## References

1. NCI-MGC;  
**National Institutes of Health, Mammalian Gene Collection (MGC)**  
<http://mgc.nci.nih.gov/>  
Unpublished.  
Position 1-1060

## Database Cross-references

**RZPD** IMAGp958C192051.  
IMAGp958C192051Q.

**UNILIB** 23347; 9630.

## Features

Key	Location	Qualifier	Value
<b>source</b>	1..1060	<b>organism</b>	Homo sapiens
		<b>lab_host</b>	DH10B (phage-resistant)
		<b>mol_type</b>	mRNA
		<b>clone_lib</b>	NIH_MGC_99
		<b>clone</b>	IMAGE:5807946
		<b>tissue_type</b>	lymphoma, cell line
		<b>note</b>	Organ: lymph; Vector: pOTB7; Site_1: XhoI; Site_2: EcoRI; cDNA made by oligo-dT priming. Directionally cloned into EcoRI/XhoI sites using the following 5' adaptor: GGCACGAG(G). Size-selected >500bp for

average insert size 1.8kb. Library constructed by Ling Hong in the laboratory of Gerald M. Rubin (University of California, Berkeley) using ZAP-cDNA synthesis kit (Stratagene) and Superscript II RT (Life Technologies). Note: this is a NIH\_MGC Library.

**db\_xref** taxon:9606

**db\_xref** RZPD:IMAGp958C192051

**db\_xref** RZPD:IMAGp958C192051Q

**db\_xref** UNILIB:23347

## Sequence

### Characteristics

**Length: 1060 BP, A Count:264, C Count:348, G Count:223, T Count:224, Others Count:1**

### Sequence

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**Go to:** General Description References Cross-references Sequence